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Minor 5'DHSR (5.1 kb subfragment) (SEQ ID NO.:6)  
-5050 AAAATATAAA AATTAGCCAG GCGTGATGTC ATGTGCCTGT AGTCCAGCT  
SP 1  
-5000 ACTCGGGAGG CTGAGGCAGG AGAACCTCTT GAATCCAGGA GGCAGAGGTT  
-4950 GCAGTGAGCA GAGATAGTGC CACTGCACTC CAGCCTGGGT GACAGAGTGA  
-4900 GACTCTGTCT CAAAAAATA AAATAAATA AAAAAATGCAG ACTGTGATTC  
-4850 AGCAGGTCTG GGTGAAGCC CAGAACTCTC TGATAAATTC AATGGCACTT  
-4800 AACTACTTGG AGGTCATGGA TGCCTTTGCT AATCTAATAG AAGCTACTGA

Major 5'DHSR (2.9 - 2.4 kb subfragments) (SEQ ID NO.:7)  
-2750 GGCTTGTGCC GAGATGTTCC CAGCACAGCC CCATGTGAGA GCTCCCTGGC  
Apa I  
-2700 TCCGGGCCCCA GTATCTGGAA TGCAGGCTCC AGCCAAATGC ATTCTCTTCT  
NFkB NFkB  
-2650 ACGGGATCTG GGAACCTCCA AAGCTGCCTC CTCAGAGTGG GAATTTCCAC  
-2600 TCACTTCTCT CACGCCAGCA CTGACCTCCC AGCGGGGGAG GGCATCTTTT  
-2550 CTTGACAGAG CAGAAGTGGG AGGCAGACAG CTGTCACTTT CCAGAAGACT  
-2500 TTCTTTTCTG ATTCATACCC TTCACCTTCC CTGTGTTTAC TGTCTGATAT  
-2450 ATGCAAAGGC CAAGTCACTT TCCAGAGATG ACAACTCCTT CCTGAAGTAG  
-2400 AGACATGCTT CCAACACTCA GAAGCCTATG TGAACACTCA GCCAGCAAAG  
SP 1  
-2350 CTGGAAGTTT TTCTCTGTA CCATGGGCTA ATTGGTCTCC TTCTCTGGAT  
AP-1  
-2300 TGTGGCTTAT CAGATAAAAA CAAGTGAGTC ATGCCACAGG ATGTCTATAA  
-2250 GCCCATTGAT TCTGGGATTC TATGAGTGAT GCTGATATGA CTAAGCCAGG  
Dra I  
-2200 AGAGACTTAT TTAAAGATCT CAGCATCTTT CAGCTTGTTA ACCTAGAGAA  
-2150 AACCCGAAGC ATGACTGGAT TATAAAGGGA AATTGAATGC GGTCCACCAA

Minor 5'DHSR (1.5 kb subfragment) (SEQ ID NO.:5)  
SP 1 Dra I  
-1300 AAGGAGGAGG CAGTGGGCTA GGAGAATCGA GAGATCAGAA TTTTAAACTC  
-1250 AGCCAGCCCA TTAACATGCC TCAAGTACTC CTATCATATT TGTAAGAGAC  
-1200 AACAGTTCAC TGAAATGAAT TCTAAGGTCT TTGGGTTTTT ATCAGTGTGC  
NF-IL 6  
-1150 TTCTGTAGTT TCTGAGGAAA TCTAAGGCAC AACTGAGGAA TGAAGTCAGG  
-1100 CTTTCCAATT CCCGAAATAC TCCTCCACTG CTTACTCATG TCCTTGGAA  
-1050 ATTAAGAAGG AAGCCAGGAG CATAGCTGCC ATAACCAGGG ATGAACCTCT

Minor 5'DHSR (0.6 kb subfragment) (SEQ ID NO.:4)  
- 500 AGGCTTCTAT GATGCTACTA TTCTGCATTT GAATGAGCAA ATGGATTTAA  
- 450 TGCATTGTCA GGGAGCCGGC CAAAGCTTGA GAGCTCCTTC CTGGCTGGGA  
- 400 GGCCCTTGG AATGTGGCCT GAAGGTAAGC TGGCAGCAG CCTGACATGC  
- 350 TTTTCATAG TTTCTCGCT TCCTTCCTTT TCTGCAGTTT TCGCTTCACA  
- 300 GAAAGCAGAA TCCTTAAAAA TAACCCTCTT AGTTCACATC TGTGGTCAGT

Minor 3'DHSR (1.3 kb subfragment) (SEQ ID NO.:3)  
+ 750 CAAAGATCAC ATTCTAGCTC TGAGGTATAG GCAGAAGCAC TGGGATTTAA  
+ 800 TGAGCTCTTT CTCTTCTCCT GCCTGCCTTT TGCTTTTTC TCATGACTCT  
+ 850 TTTCTGCTCT TAAGATCAGA ATAATCCAGT TCATCCTAAA ATGCTTTTTC

Minor 3'DHSR (2.0 kb subfragment) (SEQ ID NO.:2)  
+1550 TGCAGCTAAC TTATTTTCCC CTAGCTTTCC CCAGACACCT TGTTTTATTT  
NF-IL 6  
+1600 TATTATAATG AATTTTGTGT GTTGATGTGA AACATTATGC CTTAAGTAAT  
+1650 GTTAATTCTT ATTTAAGTTA TTGATGTTTT AAGTTTATCT TTCATGGTAC  
NF-IL 6  
+1700 TAGTGTTTTT TAGATACAGA GACTTGGGGA AATTGCTTTT CCTCTTGATC

Major 3'DHSR (2.7 - 3.4 kb subfragments) (SEQ ID NO.:1)  
NF-IL 6  
+2430 TAGGAAAATT ATAGGATCAT TAAGAAAGGA GAAGGAAGAG TGGGAGCAAA  
+2480 TACCTGGAGG TAGAAATGGT GATGATGTGT ACATCAAGCA GGGAGAAAAC  
+2530 CAATGAACCA GATGCGAATT CGGGCCCA CCAATGTCAA GGGATGACAA  
AP-1  
+2580 TTAGAAAGGA AGGTTGAGTC AAGGGATTTG AATGTTAGGG TGAAAAGTTA  
+2630 CTACTCAACT CTGTAGGTTA AAAGGAAACG TTGAGAATCT TCAGTCCAAT  
+2680 GAGGAGGGAT GTGCCATGTT TAGAGATTCA GAGATAAGTT TCAGGAAATG  
+2730 TAACCTATAG ATTTTATACA TACACAGAGA AATACGGACT AGTGAGAAGC  
+2780 TATTGCCATG GTCCAAGCAA GAGATGATGA AGGCCTAAAT ATGGAGCCAA  
+2830 AGAGGCAGCA ATGAAGAATG AGCCATGCAG GGTGAAATGC TGCATGTTGT  
+2880 AAATGGAGGA GAAAGACCTG TGAATTCAGA TATGAAAACC TCATCTTCAA  
+2930 CCCACATTTT AAGGGGGCAG CTTCCCTGAA ACCAGAATGT GTTCCCTCC  
+2980 ATTACTATAC CCCCATCCCA ATCTCAGGCA CCTGGAATCA TCCATTAA